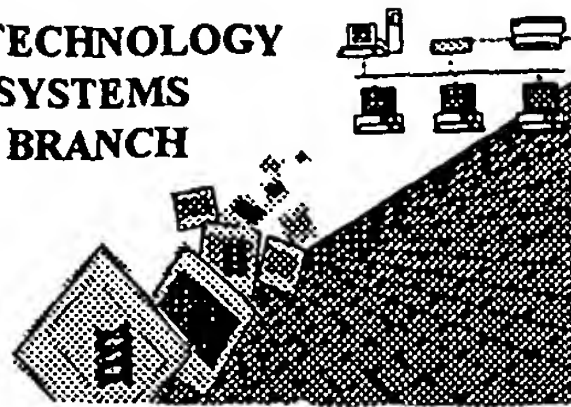


BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/053,645
Source: OIPK
Date Processed by STIC: 2/7/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

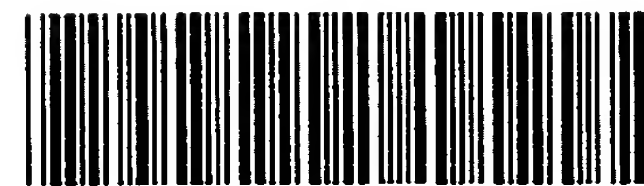
SERIAL NUMBER:

10/053,645

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220> Sequence(s) 1-19 (and more) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/053,645

DATE: 02/07/2002
TIME: 12:30:12

Input Set : A:\SEQ LISTING OF 10412-022.TXT
Output Set: N:\CRF3\02072002\J053645.raw

Does Not Comply
Corrected Diskette Needed

4 <110> APPLICANT: Robert E. Klem
6 <120> TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING A
7 CELL-PROLIFERATIVE DISORDER USING CRE DECOY OLIGOMERS, BCL-2
8 ANTISENSE OLIGOMERS, AND HYBRID OLIGOMERS THEREOF
11 <130> FILE REFERENCE: 10412-022-999
13 <140> CURRENT APPLICATION NUMBER: US/10/053,645
14 <141> CURRENT FILING DATE: 2002-01-22
16 <150> PRIOR APPLICATION NUMBER: 60/263,244
17 <151> PRIOR FILING DATE: 2001-01-22
19 <160> NUMBER OF SEQ ID NOS: 43
21 <170> SOFTWARE: FastSEQ for Windows Version 4.0
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 20
25 <212> TYPE: DNA
C--> 26 <213> ORGANISM: Artificial sequence
W--> 28 <220> FEATURE:
W--> 28 <223> OTHER INFORMATION:
28 <400> SEQUENCE: 1
29 cagcgtgcgc catccttccc
31 <210> SEQ ID NO: 2
32 <211> LENGTH: 35
33 <212> TYPE: DNA
C--> 34 <213> ORGANISM: Artificial sequence
W--> 36 <220> FEATURE:
W--> 36 <223> OTHER INFORMATION:
36 <400> SEQUENCE: 2
37 cttttcctct gggaaggatg gcgcacgctg ggaga
39 <210> SEQ ID NO: 3
40 <211> LENGTH: 20
41 <212> TYPE: DNA
C--> 42 <213> ORGANISM: Artificial sequence
W--> 44 <220> FEATURE:
W--> 44 <223> OTHER INFORMATION:
44 <400> SEQUENCE: 3
45 gatgcaccta cccagcctcc
47 <210> SEQ ID NO: 4
48 <211> LENGTH: 33
49 <212> TYPE: DNA
C--> 50 <213> ORGANISM: Artificial sequence
W--> 52 <220> FEATURE:
W--> 52 <223> OTHER INFORMATION:
52 <400> SEQUENCE: 4
53 acgggggtacg gaggtgggt aggtgcatct ggt

Artificial (global misspelling)

see item 11 on Error Summary
Sheet
(global
error)

20

35

20

33

RAW SEQUENCE LISTING

DATE: 02/07/2002

PATENT APPLICATION: US/10/053,645

TIME: 12:30:12

Input Set : A:\SEQ LISTING OF 10412-022.TXT

Output Set: N:\CRF3\02072002\J053645.raw

```

55 <210> SEQ ID NO: 5
56 <211> LENGTH: 20
57 <212> TYPE: DNA
C--> 58 <213> ORGANISM: Artificail sequence
W--> 60 <220> FEATURE:
W--> 60 <223> OTHER INFORMATION:
60 <400> SEQUENCE: 5
61 acaaaggcat cctgcagttg 20
63 <210> SEQ ID NO: 6
64 <211> LENGTH: 36
65 <212> TYPE: DNA
C--> 66 <213> ORGANISM: Artificail sequence
W--> 68 <220> FEATURE:
W--> 68 <223> OTHER INFORMATION:
68 <400> SEQUENCE: 6
69 cccccaactg caggatgcct ttgtggaact gtacgg 36
71 <210> SEQ ID NO: 7
72 <211> LENGTH: 20
73 <212> TYPE: DNA
C--> 74 <213> ORGANISM: Artificail sequence
W--> 76 <220> FEATURE:
W--> 76 <223> OTHER INFORMATION:
76 <400> SEQUENCE: 7
77 gggaaggatg gcgcacgctg 20
79 <210> SEQ ID NO: 8
80 <211> LENGTH: 17
81 <212> TYPE: DNA
C--> 82 <213> ORGANISM: Artificail sequence
W--> 84 <220> FEATURE:
W--> 84 <223> OTHER INFORMATION:
84 <400> SEQUENCE: 8
85 cgcgtgcgac cctcttg 17
87 <210> SEQ ID NO: 9
88 <211> LENGTH: 17
89 <212> TYPE: DNA
C--> 90 <213> ORGANISM: Artificail sequence
W--> 92 <220> FEATURE:
W--> 92 <223> OTHER INFORMATION:
92 <400> SEQUENCE: 9
93 taccgcgtgc gaccctc 17
95 <210> SEQ ID NO: 10
96 <211> LENGTH: 17
97 <212> TYPE: DNA
C--> 98 <213> ORGANISM: Artificail sequence
W--> 100 <220> FEATURE:
W--> 100 <223> OTHER INFORMATION:
100 <400> SEQUENCE: 10
101 tcctaccgcg tgcgacc 17
103 <210> SEQ ID NO: 11

```

RAW SEQUENCE LISTING

DATE: 02/07/2002

PATENT APPLICATION: US/10/053,645

TIME: 12:30:12

Input Set : A:\SEQ LISTING OF 10412-022.TXT

Output Set: N:\CRF3\02072002\J053645.raw

```

104 <211> LENGTH: 17
105 <212> TYPE: DNA
C--> 106 <213> ORGANISM: Artificail sequence
W--> 108 <220> FEATURE:
W--> 108 <223> OTHER INFORMATION:
108 <400> SEQUENCE: 11
109 ccttcctacc gcgtgcg 17
111 <210> SEQ ID NO: 12
112 <211> LENGTH: 17
113 <212> TYPE: DNA
C--> 114 <213> ORGANISM: Artificail sequence
W--> 116 <220> FEATURE:
W--> 116 <223> OTHER INFORMATION:
116 <400> SEQUENCE: 12
117 gacccttcct accgcgt 17
119 <210> SEQ ID NO: 13
120 <211> LENGTH: 17
121 <212> TYPE: DNA
C--> 122 <213> ORGANISM: Artificail sequence
W--> 124 <220> FEATURE:
W--> 124 <223> OTHER INFORMATION:
124 <400> SEQUENCE: 13
125 ggagaccctt cctaccg 17
127 <210> SEQ ID NO: 14
128 <211> LENGTH: 15
129 <212> TYPE: DNA
C--> 130 <213> ORGANISM: Artificail sequence
W--> 132 <220> FEATURE:
W--> 132 <223> OTHER INFORMATION:
132 <400> SEQUENCE: 14
133 gcggcggcag cgcgg 15
135 <210> SEQ ID NO: 15
136 <211> LENGTH: 15
137 <212> TYPE: DNA
C--> 138 <213> ORGANISM: Artificail sequence
W--> 140 <220> FEATURE:
W--> 140 <223> OTHER INFORMATION:
140 <400> SEQUENCE: 15
141 cggcggggcg acgga 15
143 <210> SEQ ID NO: 16
144 <211> LENGTH: 16
145 <212> TYPE: DNA
C--> 146 <213> ORGANISM: Artificail sequence
W--> 148 <220> FEATURE:
W--> 148 <223> OTHER INFORMATION:
148 <400> SEQUENCE: 16
149 cgggagcgcg gcgggc 16
151 <210> SEQ ID NO: 17
152 <211> LENGTH: 18

```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/053,645

DATE: 02/07/2002

TIME: 12:30:12

Input Set : A:\SEQ LISTING OF 10412-022.TXT

Output Set: N:\CRF3\02072002\J053645.raw

```

153 <212> TYPE: DNA
C--> 154 <213> ORGANISM: Artificail sequence
W--> 156 <220> FEATURE:
W--> 156 <223> OTHER INFORMATION:
156 <400> SEQUENCE: 17
157 tctcccagcg tgcgccat 18
159 <210> SEQ ID NO: 18
160 <211> LENGTH: 18
161 <212> TYPE: DNA
C--> 162 <213> ORGANISM: Artificail sequence
W--> 164 <220> FEATURE:
W--> 164 <223> OTHER INFORMATION:
164 <400> SEQUENCE: 18
165 tgcaatcacg ctgcgcct 18
167 <210> SEQ ID NO: 19
168 <211> LENGTH: 106
169 <212> TYPE: DNA
C--> 170 <213> ORGANISM: Artificail sequence
W--> 172 <220> FEATURE:
W--> 172 <223> OTHER INFORMATION:
172 <400> SEQUENCE: 19
173 ggcggcgccc ctccgcgcgc cctgcccgcg cgcccgcgcg gctcccgcgc gccgctctcc 60
174 ccttattggt aaaaacatgt tagaagcaat gaatgtatat aaaagc 106
176 <210> SEQ ID NO: 20
177 <211> LENGTH: 717
178 <212> TYPE: DNA
179 <213> ORGANISM: Homo Sapiens
181 <220> FEATURE:
182 <221> NAME/KEY: CDS
183 <222> LOCATION: (1)...(717)
185 <400> SEQUENCE: 20
186 atg gcg cac gct ggg aga acg ggg tac gac aac cgg gag ata gtg atg 48
187 Met Ala His Ala Gly Arg Thr Gly Tyr Asp Asn Arg Glu Ile Val Met
188 1 5 10 15
190 aag tac atc cat tat aag ctg tcg cag agg ggc tac gag tgg gat gcg 96
191 Lys Tyr Ile His Tyr Lys Leu Ser Gln Arg Gly Tyr Glu Trp Asp Ala
192 20 25 30
194 gga gat gtg ggc gcc gcg ccc ccg ggg gcc gcc ccc gca ccg ggc atc 144
195 Gly Asp Val Gly Ala Ala Pro Pro Gly Ala Ala Pro Ala Pro Gly Ile
196 35 40 45
198 ttc tcc tcc cag ccc ggg cac acg ccc cat cca gcc gca tcc cgc gac 192
199 Phe Ser Ser Gln Pro Gly His Thr Pro His Pro Ala Ala Ser Arg Asp
200 50 55 60
202 ccg gtc gcc agg acc tcg ccg ctg cag acc ccg gct gcc ccc ggc gcc 240
203 Pro Val Ala Arg Thr Ser Pro Leu Gln Thr Pro Ala Ala Pro Gly Ala
204 65 70 75 80
206 gcc gcg ggg cct gcg ctc agc ccg gtg cca cct gtg gtc cac ctg gcc 288
207 Ala Ala Gly Pro Ala Leu Ser Pro Val Pro Pro Val Val His Leu Ala
208 85 90 95

```

→ The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/053,645

DATE: 02/07/2002

TIME: 12:30:12

Input Set : A:\SEQ LISTING OF 10412-022.TXT

Output Set: N:\CRF3\02072002\J053645.raw

```

210 ctc cgc caa gcc ggc gac gac ttc tcc cgc cgc tac cgc ggc gac ttc 336
211 Leu Arg Gln Ala Gly Asp Asp Phe Ser Arg Arg Tyr Arg Gly Asp Phe
212          100          105          110
214 gcc gag atg tcc agc cag ctg cac ctg acg ccc ttc acc gcg cgg gga 384
215 Ala Glu Met Ser Ser Gln Leu His Leu Thr Pro Phe Thr Ala Arg Gly
216          115          120          125
218 cgc ttt gcc acg gtg gtg gag gag ctc ttc agg gac ggg gtg aac tgg 432
219 Arg Phe Ala Thr Val Val Glu Glu Leu Phe Arg Asp Gly Val Asn Trp
220          130          135          140
222 ggg agg att gtg gcc ttc ttt gag ttc ggt ggg gtc atg tgt gtg gag 480
223 Gly Arg Ile Val Ala Phe Phe Glu Phe Gly Gly Val Met Cys Val Glu
224 145          150          155          160
226 agc gtc aac cgg gag atg tcg ccc ctg gtg gac aac atc gcc ctg tgg 528
227 Ser Val Asn Arg Glu Met Ser Pro Leu Val Asp Asn Ile Ala Leu Trp
228          165          170          175
230 atg act gag tac ctg aac cgg cac ctg cac acc tgg atc cag gat aac 576
231 Met Thr Glu Tyr Leu Asn Arg His Leu His Thr Trp Ile Gln Asp Asn
232          180          185          190
234 gga ggc tgg gat gcc ttt gtg gaa ctg tac ggc ccc agc atg cgg cct 624
235 Gly Gly Trp Asp Ala Phe Val Glu Leu Tyr Gly Pro Ser Met Arg Pro
236          195          200          205
238 ctg ttt gat ttc tcc tgg ctg tct ctg aag act ctg ctc agt ttg gcc 672
239 Leu Phe Asp Phe Ser Trp Leu Ser Leu Lys Thr Leu Leu Ser Leu Ala
240          210          215          220
242 ctg gtg gga gct tgc atc acc ctg ggt gcc tat ctg agc cac aag 717
243 Leu Val Gly Ala Cys Ile Thr Leu Gly Ala Tyr Leu Ser His Lys
244 225          230          235
248 <210> SEQ ID NO: 21
249 <211> LENGTH: 239
250 <212> TYPE: PRT
251 <213> ORGANISM: Homo Sapiens
253 <400> SEQUENCE: 21
254 Met Ala His Ala Gly Arg Thr Gly Tyr Asp Asn Arg Glu Ile Val Met
255 1          5          10          15
256 Lys Tyr Ile His Tyr Lys Leu Ser Gln Arg Gly Tyr Glu Trp Asp Ala
257          20          25          30
258 Gly Asp Val Gly Ala Ala Pro Pro Gly Ala Ala Pro Ala Pro Gly Ile
259          35          40          45
260 Phe Ser Ser Gln Pro Gly His Thr Pro His Pro Ala Ala Ser Arg Asp
261          50          55          60
262 Pro Val Ala Arg Thr Ser Pro Leu Gln Thr Pro Ala Ala Pro Gly Ala
263 65          70          75          80
264 Ala Ala Gly Pro Ala Leu Ser Pro Val Pro Pro Val Val His Leu Ala
265          85          90          95
266 Leu Arg Gln Ala Gly Asp Asp Phe Ser Arg Arg Tyr Arg Gly Asp Phe
267          100          105          110
268 Ala Glu Met Ser Ser Gln Leu His Leu Thr Pro Phe Thr Ala Arg Gly
269          115          120          125
270 Arg Phe Ala Thr Val Val Glu Glu Leu Phe Arg Asp Gly Val Asn Trp

```

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/053,645

DATE: 02/07/2002

TIME: 12:30:13

Input Set : A:\SEQ LISTING OF 10412-022.TXT

Output Set: N:\CRF3\02072002\J053645.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application Number
L:26 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:28 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:28 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:34 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
L:36 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:36 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:42 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:44 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:44 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:50 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:52 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:52 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:58 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:60 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:60 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:66 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:68 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:68 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:74 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:76 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:76 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:82 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
L:84 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:84 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:90 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:92 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:92 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:98 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:100 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:100 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:106 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
L:108 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:108 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:114 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:116 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:116 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:122 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13
L:124 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:124 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:130 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:132 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:132 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:138 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
L:140 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:140 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:146 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16
L:148 M:258 W: Mandatory Feature missing, <220> FEATURE:

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/053,645

DATE: 02/07/2002

TIME: 12:30:13

Input Set : A:\SEQ LISTING OF 10412-022.TXT

Output Set: N:\CRF3\02072002\J053645.raw

L:148 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:154 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17
L:156 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:156 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:162 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:18
L:164 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:164 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:170 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:19
L:172 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:172 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:386 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:24
L:388 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:388 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:394 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:25
L:396 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:396 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:402 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:26
L:404 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:404 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:410 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:27
L:412 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:412 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:418 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:28
L:420 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:420 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:426 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:29
L:428 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:428 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: